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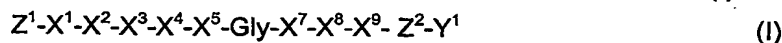
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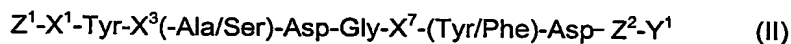
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Claims**REPLACED BY  
ART 34 AMDT**

1. A peptide comprising the amino acid sequence of formula (I)



or formula (II)



wherein

X<sup>1</sup> is an amino acid selected from the group Ser, His, Thr, Ala, Gln, Phe, Gly and Ile

X<sup>2</sup> is an amino acid selected from the group Tyr, Arg and Phe

X<sup>3</sup> is an amino acid selected from the group Tyr, Ser, Asn, Glu, Asp and Thr

X<sup>4</sup> is an amino acid selected from the group Ser, Ala, Gly, Asp and Phe

X<sup>5</sup> is an amino acid selected from the group Asp and Ser,

X<sup>7</sup> is an amino acid selected from the group Thr, Val, Met, Ser, Trp, Tyr, Leu and Ala

X<sup>8</sup> is an amino acid selected from the group Tyr, Phe and Leu

X<sup>9</sup> is an amino acid selected from the group Asp, Ser and Glu

Z<sup>1</sup> represent an amino acid residue capable of forming a disulphide bond, preferably a cysteine or a homocysteine residue, or a residue capable of forming a thioether preferably the residue is Q-C(=O) wherein Q represents  $-(CH_2)_n$  or  $-(CH_2)_n-C_6H_4$  where n represents a positive integer 1 to 10 or is absent and

Z<sup>2</sup> represent an amino acid residue capable of forming a disulphide bond, preferably a cysteine or a homocysteine residue or is absent

Y<sup>1</sup> represents 1-10 amino acids or is absent

or pharmaceutically acceptable salts thereof.

2. A peptide according to claim 1 of the amino acid sequence

Cys-Ser-Tyr-Tyr-Ser-Asp-Gly-Val-Tyr-Asp-Cys, (SEQ ID NO 1),

Cys-His-Tyr-Ser-Ser-Asp-Gly-Thr-Tyr-Asp-Cys, (SEQ ID NO 2),

Cys-Thr-Tyr-Asn-Gly-Asp-Gly-Ser-Phe-Asp-Cys, (SEQ ID NO 3),

Cys-Ala-Tyr-Glu-Ala-Asp-Gly-Trp-Phe-Asp-Cys, (SEQ ID NO 4),

Cys-Ser-Tyr-Ser-Ala-Asp-Gly-Thr-Leu-Asp-Cys, (SEQ ID NO 5),

Cys-Gln-Tyr-Asp-Ser-Ser-Gly-Met-Tyr-Asp-Cys, (SEQ ID NO 6),

Cys-Phe-Phe-Asp-Ser-Ser-Gly-Tyr-Phe-Asp-Cys, (SEQ ID NO 7),

Cys-Thr-Tyr-Ser-Ala-Asp-Gly-Leu-Tyr-Asp-Cys, (SEQ ID NO 8),

Cys-His-Phe-Asp-Gly-Asp-Gly-Ser-Tyr-Asp-Cys, (SEQ ID NO 9),